

#3

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/430,806A

DATE: 10/20/2000
TIME: 12:18:21

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4 110> APPLICANT: Vinkemeier, Uwe
6 Barnell Jr., James E.
10 120> TITLE OF INVENTION: PURIFIED STAT PROTEINS AND METHODS OF PURIFYING THEREOF
14 130> FILE REFERENCE: 600-1-182 N
17 <110> CURRENT APPLICATION NUMBER: 09/430,806A
18 <141> CURRENT FILING DATE: 1996-11-02
21 <150> PRIOR APPLICATION NUMBER: 08/351,130
23 <151> PRIOR FILING DATE: 1993-10-15
27 <150> PRIOR APPLICATION NUMBER: 60/028,176
29 <151> PRIOR FILING DATE: 1996-10-15
33 <160> NUMBER OF SEQ ID NOS: 16
37 <170> SOFTWARE: Patentin Ver. 2.0
41 <210> SEQ ID NO: 1
43 <211> LENGTH: 750
45 <212> TYPE: PRT
47 <213> ORGANISM: Homo sapiens
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61 20 25 30
65 Tyr Leu Ala Gln Trp Leu Glu Lys Gln Asp Trp Glu His Ala Ala Asn
67 35 40 45
71 Asp Val Ser Phe Ala Thr Ile Arg Phe His Asp Leu Leu Ser Gln Leu
73 40 45 50 55 60
77 Asp Asp Gln Tyr Ser Arg Phe Ser Leu Glu Asn Asn Phe Leu Leu Gln
79 65 70 75 80
83 His Asn Ile Arg Lys Ser Lys Arg Asn Leu Gln Asp Asn Phe Gln Glu
85 80 85 90 95
89 Asp Pro Ile Gln Met Ser Met Ile Ile Tyr Ser Cys Leu Lys Glu Glu
91 100 105 110
95 Arg Lys Ile Leu Glu Asn Ala Gln Arg Phe Asn Gln Ala Gln Ser Gly
97 115 120 125
101 Asn Ile Gln Ser Thr Val Met Leu Asp Lys Gln Lys Glu Leu Asp Ser
103 130 135 140 145
107 Lys Val Arg Asn Val Lys Asp Lys Val Met Cys Ile Glu His Glu Ile
109 145 150 155 160
113 Lys Ser Leu Glu Asp Leu Gln Asp Glu Tyr Asp Phe Lys Cys Lys Thr
115 165 170 175
119 Leu Gln Asn Arg Glu His Glu Thr Asn Glu Val Ala Lys Ser Asp Gln
121 180 185 190
125 Lys Gln Glu Gln Leu Leu Leu Lys Lys Met Tyr Leu Met Leu Asp Asn
127 195 200 205
131 Lys Arg Lys Glu Val Val His Ile Ile Glu Leu Leu Asn Val Thr
133 210 215 220
137 Glu Leu Thr Gln Asn Ala Leu Ile Asn Asp Glu Leu Val Glu Trp Lys
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149 225

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151 Val Arg Gln Gln Ile Lys Lys Ile Gln Gln Lys Tyr Thr
152 275 280 285
153 Tyr Glu His Asp Pro Ile Thr Lys Asn Lys Gln Val Leu Trp Asp Arg
161 290 295 300
163 Thr Phe Ser Leu Phe Gln Gln Ile Gln Ser Ser Phe Val Val Glu
167 305 310 315 320
169 325 330 335
172 Arg Gln Pro Cys Met Pro Thr His Pro Gln Arg Pro Leu Val Leu Lys
173 340 345 350
176 Thr Gly Val Gln Phe Thr Val Lys Leu Arg Leu Leu Val Lys Leu Gln
177 355 360 365
181 Glu Leu Asn Tyr Asn Leu Lys Val Lys Val Leu Phe Asp Lys Asp Val
185 370 375 380
187 Asn Glu Arg Asn Thr Val Lys Gly Phe Arg Lys Phe Asn Ile Leu Gly
191 385 390 395 400
192 Thr His Thr Lys Val Met Asn Met Glu Glu Ser Thr Asn Gly Ser Leu
193 405 410 415 420
197 425 430 435 440 445
203 Ala Ala Glu Phe Arg His Ile Gln Leu Lys Glu Gln Lys Asn Ala Gly
205 450 455 460 465 470 475 480
209 Thr Arg Thr Asn Glu Gly Pro Leu Ile Val Thr Glu Glu Leu His Ser
211 485 490 495 500 505 510 515
215 Leu Ser Phe Glu Thr Gln Ile Cys Gln Pro Gly Leu Val Ile Asp Leu
217 520 525 530 535 540 545 550
221 Glu Thr Thr Ser Leu Pro Val Val Val Ile Ser Asn Val Ser Gln Leu
223 555 560 565 570 575 580 585
227 Pro Ser Gly Trp Ala Ser Ile Leu Ile Tyr Asn Met Leu Val Ala Glu
229 590 595 600 605 610 615 620
233 625 630 635 640 645 650 655
235 660 665 670 675 680 685 690
239 Gln Leu Ser Glu Val Leu Ser Trp Gln Phe Ser Ser Val Thr Lys Arg
241 695 700 705 710 715 720 725
245 Gly Leu Asn Val Asp Gln Leu Asn Met Leu Gly Glu Lys Leu Leu Gly
247 730 735 740 745 750 755 760
251 Pro Asn Ala Ser Pro Asp Gly Ile Ile Pro Trp Thr Arg Phe Cys Lys
253 765 770 775 780 785 790 795
257 Glu Asn Ile Asn Asp Lys Asn Phe Pro Phe Trp Leu Ile Glu Ser
259 800 805 810 815 820 825 830
263 Ile Leu Glu Leu Ile Lys Lys His Ile Leu Pro Leu Trp Asn Asp Gly
265 835 840 845 850 855 860 865
269 Cys Ile Met Gly Phe Ile Ser Lys Glu Arg Glu Arg Ala Leu Leu Lys
271 870 875 880 885 890 895 900
275 Asp Gln Gln Pro Gly Thr Phe Leu Arg Phe Ser Glu Ser Ser Arg
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281 Glu Gly Ala Ile Thr Phe Thr Ile Val Glu Arg Ser Gln Asn Gly Gly
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287 Glu Pro Asp Phe His Ala Val Glu Pro Tyr Thr Lys Lys Glu Leu Ser

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296 Glu Asn Ile Pro Pro Glu Asn Pro Leu Lys Tyr Leu Tyr Pro Asn Ile Asp
301 660 665 670
305 Lys Asp His Ala Phe Gly Lys Tyr Tyr Ser Arg Pro Lys Glu Ala Pro
307 675 680 685
311 Glu Pro Met Glu Leu Asp Gly Pro Lys Gly Ile Lys Thr
314 690 695 700
315 Glu Leu Ile Ser Val Ser Glu Val His Pro Ser Arg Leu Gln Thr Ile
319 705 710 715 720
323 Asp Asn Leu Leu Pro Met Ser Pro Glu Glu Phe Asp Glu Val Ser Arg
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329 740 745 750
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334 310 - SEQ ID NO: 2
334 311 - LENGTH: 712
349 312 - TYPE: PRT
342 313 - ORGANISM: Homo sapiens
346 3400 - SEQUENCE: 2
348 Met Ser Glu Trp Tyr Glu Leu Gln Gln Leu Asp Ser Lys Phe Leu Gln
353 1 5 10 15
350 20 25 30
354 Gln Val His Gln Leu Tyr Asp Asp Ser Phe Pro Met Glu Ile Arg Gln
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350 Tyr Leu Ala Gln Trp Leu Gln Lys Gln Asp Trp Glu His Ala Ala Asn
362 35 40 45
366 Asp Val Ser Phe Ala Thr Ile Arg Phe His Asp Leu Leu Ser Gln Leu
368 50 55 60
372 Asp Asp Gln Tyr Ser Arg Phe Ser Leu Glu Asn Asn Phe Leu Leu Gln
374 65 70 75 80
378 His Asn Ile Arg Lys Ser Lys Arg Asn Leu Gln Asp Asn Phe Gln Gln
380 85 90 95
384 Asp Pro Ile Gln Met Ser Met Ile Ile Tyr Ser Cys Leu Lys Glu Glu
386 100 105 110
390 Arg Lys Ile Leu Glu Asn Ala Gln Arg Phe Asn Gln Ala Gln Ser Gln
392 115 120 125
396 Asn Ile Gln Ser Thr Val Met Leu Asp Lys Gln Lys Glu Leu Asp Ser
408 130 135 140
402 Lys Val Arg Asn Val Lys Asp Lys Val Met Cys Ile Glu His Gln Ile
404 143 150 155 160
408 Lys Ser Leu Glu Asp Leu Gln Asp Glu Tyr Asp Phe Lys Cys Lys Thr
410 165 170 175
414 Leu Gln Asn Arg Glu His Glu Thr Asn Gly Val Ala Lys Ser Asp Gln
416 180 185 190
420 Lys Gln Gln Gln Leu Leu Lys Lys Met Tyr Leu Met Leu Asp Asn
422 195 200 205
426 Lys Arg Lys Glu Val Val His Lys Ile Ile Glu Leu Leu Asn Val Thr
428 210 215 220
432 Glu Leu Thr Gln Asn Ala Leu Ile Asn Asp Glu Leu Val Glu Trp Lys

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516	Glu Thr Thr Ser Leu Pro Val Val Val Ile Ser Asn Val Ser Gln Leu			
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VERIFICATION SUMMARY

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